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1	<u>CLAIMS</u>
2	What is claimed is:
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4	Claim 1. A biopolymer marker selected from the group
5	consisting of sequence ID IHWESASLLR,
6	<pre>ILLQGTPVAQMTEDAVDAER, (K) DFDFVPPVVR(W) or at least one</pre>
7	analyte thereof useful in indicating at least one
8	particular disease state.
9	
10	Claim 2. The biopolymer marker of claim 1 wherein
11	said disease state is predictive of Type II diabetes.
12	
13	Claim 3. A method for evidencing and categorizing at
14	least one disease state comprising:
15	obtaining a sample from a patient;
16	conducting mass spectrometric analysis on said
17	sample;
18	evidencing and categorizing at least one biopolymer
19	marker sequence or analyte thereof isolated from said
20	sample; and,
21	comparing said at least one isolated biopolymer
22	marker sequence or analyte thereof to the biopolymer
23	marker sequence as set forth in claim 1;
24	wherein correlation of said isolated biopolymer

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1	marker and said biopolymer marker sequence as set forth in
2	claim 1 evidences and categorizes said at least one
3	disease state.
4	
5	Claim 4. The method of claim 3, wherein said step
6	of evidencing and categorizing is particularly directed to
7	biopolymer markers or analytes thereof linked to at least
8	one risk of disease development of said patient.
9	
10	Claim 5. The method of claim 3, wherein said step
11	of evidencing and categorizing is particularly directed to
12	biopolymer markers or analytes thereof related to the
13	existence of a particular disease state.
14	
15	Claim 6. The method of claim 3, wherein the sample
16	is an unfractionated body fluid or a tissue sample.
17	
18	
19	Claim 7. The method of claim 3, wherein said sample
20	is at least one of the group consisting of blood, blood
21	products, urine, saliva, cerebrospinal fluid, and lymph.
22	
23	Claim 8. The method of claim 3, wherein said mass
24	spectrometric analysis is selected from the group

1 consisting of Surface Enhanced Laser Desorption Ionization 2 (SELDI) mass spectrometry (MS), Maldi Qq TOF, MS/MS, 3 TOF-TOF, and ESI-Q-TOF or an ION-TRAP. 4 5 Claim 9. The method of claim 3, wherein said 6 patient is a human. 7 8 A diagnostic assay kit for determining Claim 10. 9 the presence of the biopolymer marker or analyte thereof 10 of claim 1 comprising: 11 at least one biochemical material which is capable of specifically binding with a biomolecule which includes at 12 least said biopolymer marker or analyte thereof, and 13 14 means for determining binding between said 15 U D 16 biochemical material and said biomolecule: whereby at least one analysis to determine a presence 17 of a marker, analyte thereof, or a biochemical material 18 specific thereto, is carried out on a sample. 19 20 Claim 11. The diagnostic assay kit of claim 10, 21 wherein said biochemical material or biomolecule is 22 immobilized on a solid support. 23 24 Claim 12. The diagnostic assay kit of claim 10

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1	including:
2	at least one labeled biochemical material.
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4	Claim 13. The diagnostic assay kit of claim 10,
5	wherein said biochemical material is an antibody.
6	
7	Claim 14. The diagnostic assay kit of claim 12,
8	wherein said labeled biochemical material is an antibody.
9	
10	Claim 15. The diagnostic assay kit of claim 10,
11	wherein the sample is an unfractionated body fluid or a
12	tissue sample.
13	
14	Claim 16. The diagnostic assay kit of claim 10,
15	wherein said sample is at least one of the group
16	consisting of blood, blood products, urine, saliva,
17	cerebrospinal fluid, and lymph.
18	
19	Claim 17. The diagnostic assay kit of claim 10,
20	wherein said biochemical material is at least one
21	monoclonal antibody specific therefore.
22	
23	Claim 18. A kit for diagnosing, determining risk-
24	assessment, and identifying therapeutic avenues related to

The kit of claim 20, wherein said labeled

Claim 22.

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a disease state comprising:

1 biochemical material is an antibody. 2 The kit of claim 18, wherein the sample is 3 Claim 23. an unfractionated body fluid or a tissue sample. 4 5 The kit of claim 18, wherein said sample 6 Claim 24. is at least one of the group consisting of blood, blood 7 products, urine, saliva, cerebrospinal fluid, and lymph. 8 9 □ 10 Claim 25. The kit of claim 18, wherein said 10 11 11 12 11 12 biochemical material is at least one monoclonal antibody specific therefore. □ 13 ∀ The kit of claim 18, wherein said 14 Claim 26. H 15 diagnosing, determining risk assessment, and identifying 16 therapeutic avenues is carried out on a single sample. 17 The kit of claim 18, wherein said 18 Claim 27.

diagnosing, determining risk assessment, and identifying therapeutic avenues is carried out on multiple samples such that at least one analysis is carried out on a first sample and at least another analysis is carried out on a second sample.

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1	Claim 28. The kit of claim 27, wherein said first
2	and second samples are obtained at different time periods.
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4	Claim 29. Polyclonal antibodies produced against a
5	marker sequence ID selected from the group consisting of
6	sequence ID IHWESASLLR, ILLQGTPVAQMTEDAVDAER,
7	(K)DFDFVPPVVR(W) or at least one analyte thereof in at
8	least one animal host.
9	
10	Claim 30. An antibody that specifically binds a
11	biopolymer including a marker selected from the group
12	consisting of sequence ID IHWESASLLR,
13	<pre>ILLQGTPVAQMTEDAVDAER, (K)DFDFVPPVVR(W) or at least one</pre>
14	analyte thereof.
15	
16	Claim 31. The antibody of claim 30 that is a
17	monoclonal antibody.
18	
19	Claim 32. The antibody of claim 30 that is a
20	polyclonal antibody.
21	
22	Claim 33. A process for identifying therapeutic
23	avenues related to a disease state comprising:
24	conducting an analysis as provided by the kit of

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1 claim 18; and interacting with a biopolymer selected from the group 2 3 consisting of sequence ID IHWESASLLR, ILLQGTPVAQMTEDAVDAER, (K)DFDFVPPVVR(W) or at least one 4 analyte thereof; 5 6 whereby therapeutic avenues are developed. 7 The process for identifying therapeutic 8 Claim 34. 9 avenues related to a disease state in accordance with 10 claim 33, wherein said therapeutic avenues regulate the 11 presence or absence of the biopolymer selected from the 12 group consisting of sequence ID IHWESASLLR, ILLQGTPVAQMTEDAVDAER, (K) DFDFVPPVVR(W) or at least one analyte thereof. 15 The process for identifying therapeutic Claim 35. 16 avenues related to a disease state in accordance with 17 18 claim 33, wherein said therapeutic avenues developed 19 include at least one avenue selected from a group 20 consisting of 1)utilization and recognition of said biopolymer markers, variants or moieties thereof as direct 21 22 therapeutic modalities, either alone or in conjunction

with an effective amount of a pharmaceutically effective

carrier; 2) validation of therapeutic modalities or disease

1 preventative agents as a function of biopolymer marker 2 presence or concentration; 3) treatment or prevention of a 3 disease state by formation of disease intervention 4 modalities; 4) use of biopolymer markers or moieties thereof as a means of elucidating therapeutically viable 5 6 agents, 5) instigation of a therapeutic immunological response; and 6) synthesis of molecular structures related 7 to said biopolymer markers, moieties or variants thereof 9 which are constructed and arranged to therapeutically intervene in said disease state. Claim 36. The process for identifying therapeutic avenues related to a disease state in accordance with 13 claim 35, wherein said treatment or prevention of a disease state by formation of disease intervention modalities is the formation of biopolymer/ligand 16 17 conjugates which intervene at receptor sites to prevent, 18 delay or reverse a disease process. 19 20 Claim 37. The process for identifying therapeutic avenues related to a disease state in accordance with

- 21
- 22 claim 35, wherein said means of elucidating
- 23 therapeutically viable agents includes use of a
- bacteriophage peptide display library or a bacteriophage 24

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antibody library.

Claim 38. A process for regulating a disease state by controlling the presence or absence of a biopolymer selected from the group consisting of sequence ID

HWESASLLR, ILLQGTPVAQMTEDAVDAER, (K)DFDFVPPVVR(W) or at least one analyte thereof.
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